SEQUENCE LISTING

Witte, Owen N.
Saffran, Douglas C.
Jakobovits, Aya

<120> PSCA: PROSTATE STEM CELL ANTIGEN AND USES THEREOF

<130> 30435.54USD3

<140> 09/934,773

<141> 2001-08-21

<150> 09/564,329

<151> 2000-05**-**03

<150> 09/359,326\

<151> 1999-07-20

<150> 09/318,503

<151> 1999-05-25

<150> 09/251,835

<151> 1999-02-17

<150> 09/203,939

<151> 1998-12-02

<150> 09/038,261

<151> 1998-03-10

<150> 60/124,658

<151> 1999-03-16

<150> 60/120,536

<151> 1999-02-17

<150> 60/113,230

<151> 1998-12-21

<150> 60/074,675

<151> 1998-02-13

<150> 60/071,141

<151> 1998-01-12

<150> 60/228,816

<151> 1997-03-10

<160> 27

<170> PatentIn Ver. 2.0

<210>.1

<211> 998

<212> DNA

Amos -

REPLACED BY

```
<213> Homo sapiens
   <220>
   <221> misc feature
   <222> (543)
   <223> any nucleotide (i.e., a, c, g or t)
   <220>
   <221> misc_feature
   <222> (580)
   <223> any nucleotide (i.e., a, c, g or t)
   <220>
   <221> misc_feature
   <222> (584)
  <223> any nucleotide (i.e., a, c, g or t)
   <220>
   <221> misc feature
   <222> (604)
   <223> any nucleotide (i.e., a, c, g or t)
   <220>
   <221> misc_feature
   <222> (608)
   <223> any nucleotide (i.e., a, c, g or t)
  <220>
   <221> misc_feature
  <222> (615)
   <223> any nucleotide (i.e., a, c, g or t)
  <220>
  <221> misc_feature 🤌
  <222> (636)
  <223> any nucleotide (i.e., a, c, g or t)
  <220>
  <221> misc feature
  <222> (640).
  <223> any nucleotide (i.e., a, c, g or t)
  <220>
  <221> misc_feature
  <222> (646)
<223> any nucleotide (i.e., a, c, g or t)
  <220>
  <221> misc_feature
  <222> (697)
  <223> any nucleotide (i.e., a, c, g or t)
 ..<220>
  <221> misc feature
  <222> (926)
  <223> any nucleotide (i.e., a, c, g or t)
```

```
agggagagge agtgaccatg aaggetgtge tgettgeeet gttgatggea ggettggeee 60
 tgcagccagg cactgccctg ctgtgctact cctgcaaagc ccaggtgagc aacgaggact 120
 gcctgcaggt ggagaactgc acccagctgg gggagcagtg ctggaccgcg cgcatccgcg 180
 cagttggcct cctgaccgtc atcagcaaag gctgcagctt gaactgcgtg gatgactcac 240
 aggactacta cgtgggcaag aagaacatca cgtgctgtga caccgacttg tgcaacgcca 300
 geggggccca tgccctgcag ccggctgccg ccatccttgc gctgctccct gcactcggcc 360
 tgctgctctg gggacccggc cagctatagg ctctgggggg ccccgctgca gcccacactg 420
 ggtgtggtgc cccaggcctt tgtgccactc ctcacagaac ctggcccagt gggagcctgt 480
 cctggttcct gaggcacatc ctaacgcaag tttgaccatg tatgtttgca ccccttttcc 540
 ccnaaccetg acctteceat gggeetttte caggatteen acenggeaga teagttttag 600
 tganacanat ccgcntgcag atggcccctc caaccntttn tgttgntgtt tccatggccc 660
 agcattttcc accettaace etgtgttcag gcacttnttc ccccaggaag cettcectge 720
 ccaccccatt tatgaattga gccaggtttg gtccgtggtg tcccccgcac ccagcagggg 780
 acaggcaatc aggagggccc agtaaaggct gagatgaagt ggactgagta gaactggagg 840
 acaagagttg acgtgagttc ctgggagttt ccagagatgg ggcctggagg cctggaggaa 900
 ggggccaggc ctcacatttg tggggntccc gaatggcagc ctgagcacag cgtaggccct 960
 taataaacac ctgttggata agccaaaaaa aaaaaaaa
<210> 2
 <211> 123
 <212> PRT
 <213> Homo sapiens
<220>
<221> PEPTIDE
<222> (50)..(64)
<220>
<221> PEPTIDE
<222> (71) .. (82)
<220>
<221> PEPTIDE
<222> (67)..(81)
<400> 2
Met Lys Ala Val Leu Leu Ala Leu Leu Met Ala Gly Leu Ala Leu Gln
Pro Gly Thr Ala Leu Leu Cys Tyr Ser Cys Lys Ala Gln Val Ser Asn
             20
Glu Asp Cys Leu Gln Val Glu Asn Cys Thr Gln Leu Gly Glu Gln Cys
                             40.
Trp Thr Ala Arg Ile Arg Ala Val Gly Leu Leu Thr Val Ile Ser Lys
     50
Gly Cys Ser Leu Asn Cys Val Asp Asp Ser Gln Asp Tyr Tyr Val Gly
Lys Lys Asn Ile Thr Cys Cys Asp Thr Asp Leu Cys Asn Ala Ser Gly
Ala His Ala Leu Gln Pro Ala Ala Ala Ile Leu Ala Leu Leu Pro Ala
            100
                                105
                                                    110
```

<400> 1

```
Leu Gly Leu Leu Trp Gly Pro Gly Gln Leu
        · 115
  <210> 3
  <211> 441
  <212> DNA
  <213> Mus musculus
 <400> 3
 atgaagacag tttttttat cctgctggcc acctacttag ccctqcatcc aggtgctgct 60
 ctgcagtgct attcatgcac agcacagatg aacaacagag actgtctgaa tgtacagaac 120
tgcagcctgg accagcacag ttgctttaca tcgcgcatcc gggccattgg actcgtgaca 180
 gttatcagta agggctgcag ctcacagtgt gaggatgact cggagaacta ctatttgggc 240
 aagaagaaca tcacgtgctg ctactctgac ctgtgcaatg tcaacggggc ccacaccctg 300
 aagccaccca ccaccctggg gctgctgacc gtgctctgca gcctgttgct gtggggctcc 360
 agccgtctgt aggctctggg agagcctacc atagcccgat tgtgaaggga tgagctgcac 420
 tccaccccac ccccacacag g
 <210> 4
 <211> 123
 <212> PRT
 <213> Mus musculus
 <400> 4
 Met Lys Thr Val Phe Phe Ile Leu Leu Ala Thr Tyr Leu Ala Leu His
 Pro Gly Ala Ala Leu Gln Cys Tyr Ser Cys Thr Ala Gln Met Asn Asn
              20
 Arg Asp Cys Leu Asn Val Gln Asn Cys Ser Leu Asp Gln His Ser Cys
 Phe Thr Ser Arg Ile Arg Ala Ile Gly Leu Val Thr Val Ile Ser Lys
                          55
 Gly Cys Ser Ser Gln Cys Glu Asp Asp Ser Glu Asn Tyr Tyr Leu Gly
  65
                                           75
 Lys Lys Asn Ile Thr Cys Cys Tyr Ser Asp Leu Cys Asn Val Asn Gly
                                      90 -
 Ala His Thr Leu Lys Pro Pro Thr Thr Leu Gly Leu Leu Thr Val Leu
             100 ·
                                 105
 Cys Ser Leu Leu Trp Gly Ser Ser Arg Leu
```

<210> 5 <211> 131 <212> PRT <213> Homo sapiens

<400> 5

Met Lys Ile Phe Leu Pro Val Leu Leu Ala Ala Leu Leu Gly Val Glu 1 5 15

Arg Ala Ser Ser Leu Met Cys Phe Ser Cys Leu Asn Gln Lys Ser Asn 20 25 30

Leu Tyr Cys Leu Lys Pro Thr Ile Cys Ser Asp Gln Asp Asn Tyr Cys
35 40 45

Val Thr Val Ser Ala Ser Ala Gly Ile Gly Asn Leu Val Thr Phe Gly 50 60

His Ser Leu Ser Lys Thr Cys Ser Pro Ala Cys Pro Ile Pro Glu Gly 65 70 75 80

Val Asn Val Gly Val Ala Ser Met Gly Ile Ser Cys Cys Gln Ser Phe
85 90 95

Leu Cys Asn Phe Ser Ala Ala Asp Gly Gly Leu Arg Ala Ser Val Thr 100 105 110

Leu Leu Gly Ala Gly Leu Leu Leu Ser Leu Leu Pro Ala Leu Leu Arg 115 120 125

Phe Gly Pro 130

<210> 6

<211> 123

<212> PRT

<213> Homo sapiens

<400> 6

Met Lys Ala Val Leu Leu Ala Leu Leu Met Ala Gly Leu Ala Leu Gln
1 5 10 15

Pro Gly Thr Ala Leu Leu Cys Tyr Ser Cys Lys Ala Gln Val Ser Asn 20 25 30

Glu Asp Cys Leu Gln Val Glu Asn Cys Thr Gln Leu Gly Glu Gln Cys 35 40 45

Trp Thr Ala Arg Ile Arg Ala Val Gly Leu Leu Thr Val Ile Ser Lys
50 55 60

Gly Cys Ser Leu Asn Cys Val Asp Asp Ser Gln Asp Tyr Tyr Val Gly
65 70 ...75 80

Lys Lys Asn Ile Thr Cys Cys Asp Thr Asp Leu Cys Asn Ala Ser Gly
85 90 95

Ala His Ala Leu Gln Pro Ala Ala Ala Ile Leu Ala Leu Leu Pro Ala 100 105 110

Leu Gly Leu Leu Eu Trp Gly Pro Gly Gln Leu
115 120

```
<210> 7
 <211> 123
 <212> PRT
 <213> Mus musculus
 <400> 7
 Met Lys Thr Val Leu Phe Leu Leu Leu Ala Thr Tyr Leu Ala Leu His
 Pro Gly Ala Ala Leu Gln Cys Tyr Ser Cys Thr Ala Gln Met Asn Asn
                                - 25
 Arg Asp Cys Leu Asn Val Gln Asn Cys Ser Leu Asp Gln His Ser Cys
 Phe Thr Ser Arg Ile Arg Ala Ile Gly Leu Val Thr Val Ile Ser Lys
 Gly Cys Ser Ser Gln Cys Glu Asp Asp Ser Glu Asn Tyr Tyr Leu Gly
                      70
                                          75
 Lys Lys Asn Ile Thr Cys Cys Tyr Ser Asp Leu Cys Asn Val Asn Gly
                                     90
 Ala His Thr Leu Lys Pro Pro Thr Thr Leu Gly Leu Leu Thr Val Leu
 Cys Ser Leu Leu Trp Gly Ser Ser Arg Leu
 <210> 8
 <211> 20
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: RT-PCR PRIMER
 <400> 8
 ttctcctgct ggccacctac
                                                                    20
 <210> 9
 <211> 20
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: RT-PCR PRIMER
· <400> 9
 gcagctcatc ccttcacaat
                                                                    20
<210> 10
```

<211> 408

```
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: MONOCLONAL
      ANTIBODY 1G8
<400> 10
tgcttcttcc, tgatggcagt ggttatagga gtcaattcag aggttcagct gcagcagtct 60
ggggcagaac ttgtgaggtc aggggcctca gtcaagttgt cctgcacagc ttctggcttc 120
aacattaaag actactatat acactgggtg aatcagaggc ctgaccaggg cctggagtgg 180
attggatgga ttgatcctga gaatggtgac actgaatttg tcccgaagtt ccagggcaag 240
gecaetatga etgeagaeat ttteteeaae acageetaee tgeaeeteag eageetgaea 300
tetgaagaca etgeegteta ttaetgtaaa aeggggggtt tetggggeea agggaetetg 360
gtcactgtct ctgcagccaa aacgacaccc ccatctgtct atccactg
<210> 11
<211> 136
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: MONOCLONAL
      ANTIBODY 1G8
<400> 11
Cys Phe Phe Leu Met Ala Val Val Ile Gly Val Asn Ser Glu Val Gln
Leu Gln Gln Ser Gly Ala Glu Leu Val Arg Ser Gly Ala Ser Val Lys
            20
Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Tyr Tyr Ile His
Trp Val Asn Gln Arg Pro Asp Gln Gly Leu Glu Trp Ile Gly Trp Ile
                         55
Asp Pro Glu Asn Gly Asp Thr Glu Phe Val Pro Lys Phe Gln Gly Lys
Ala Thr Met Thr Ala Asp Ile Phe Ser Asn Thr Ala Tyr Leu His Leu
                                     90
Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys Lys Thr Gly
            100
                                105
                                                    110
Gly Phe Trp Gly Gln Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr
                                                125
Thr Pro Pro Ser Val Tyr Pro Leu
   130
<210> 12-
```

<212> DNA

<211> 426 <212> DNA J.

```
<213> Artificial Sequence
 <223> Description of Artificial Sequence: MONOCLONAL
      ANTIBODY 4A10
 <400> 12
ttggtagcaa cagcctcaga tgtccactcc caggtccaac tgcagcaacc tgggtctgaa 60
ctggtgaggc ctggaacttc agtgaagctg tcctgcaagg cttctggcta tacattctcc 120
agctactgga tgcactgggt gaagcagagg cctggacaag gccttgagtg gattggaaat 180
attgaccetg gtagtggtta cactaactac getgagaace teaagaccaa ggecacaetg 240
actgtagaca catectecag cacagectae atgeagetea geageetgae atetgaggae 300
tctgcagtct attactgtac aagccgatct actatgatta cgacgggatt tgcttactgg 360
ggccaaggga ctctggtcac tgtctctgca gctacaacaa cagccccatc tgtctatcca 420
ctggcc
<210> 13
<211> 142
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: MONOCLONAL
      ANTIBODY 4A10
<400> 13
Leu Val Ala Thr Ala Ser Asp Val His Ser Gln Val Gln Leu Gln Gln
Pro Gly Ser Glu Leu Val Arg Pro Gly Thr Ser Val Lys Leu Ser Cys
             20
Lys Ala Ser Gly Tyr Thr Phe Ser Ser Tyr Trp Met His Trp Val Lys
Gln Arg Pro Gly Gln Gly Leu Glu Trp Ile Gly Asn Ile Asp Pro Gly
                         55
Ser Gly Tyr Thr Asn Tyr Ala Glu Asn Leu Lys Thr Lys Ala Thr Leu
Thr Val Asp Thr Ser Ser Ser Thr Ala Tyr Met Gln Leu Ser Ser Leu
                                     90
Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Thr Ser Arg Ser Thr Met
            100
                               105
                                                    110
Ile Thr Thr Gly Phe Ala Tyr Trp Gly Gln Gly Thr Leu Val Thr Val
Ser Ala Ala Thr Thr Thr Ala Pro Ser Val Tyr Pro Leu Ala
    130
                        135
                                            140
```

<210> 14 <211> 453

<212> DNA

```
<213> Artificial Sequence
<223> Description of Artificial Sequence: MONOCLONAL
      ANTIBODY 2H9
<400> 14
aatgacttcg ggttgagctg ggtttttatt attgttcttt taaaaggggt ccggagtgaa 60
gtgaggettg aggagtetgg aggaggetgg gtgeaacetg gaggateeat gaaactetee 120
tgtgtagcct ctggatttac tttcagtaat tactggatga cttgggtccg ccagtctcca 180
gagaaggggc ttgagtgggt tgctgaaatt cgattgagat ctgaaaatta tgcaacacat 240
tatgcggagt ctgtgaaagg gaaattcacc atctcaagag atgattccag aagtcgtctc 300
tacctgcaaa tgaacaactt aagacctgaa gacagtggaa tttattactg tacagatggt 360
ctgggacgac ctaactgggg ccaagggact ctggtcactg tctctgcagc caaaacgaca 420
cccccatctg tctatccact ggccccttgt gta
<210> 15
<211> 151
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: MONOCLONAL
      ANTIBODY 2H9
<400> 15
Asn Asp Phe Gly Leu Ser Trp Val Phe Ile Ile Val Leu Leu Lys Gly
                                     10
Val Arg Ser Glu Val Arg Leu Glu Glu Ser Gly Gly Gly Trp Val Gln
             20 .
                                 25
Pro Gly Gly Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Thr Phe
                             40
Ser Asn Tyr Trp Met Thr Trp Val Arg Gln Ser Pro Glu Lys Gly Leu
                        55
                                             . 60
Glu Trp Val Ala Glu Ile Arg Leu Arg Ser Glu Asn Tyr Ala Thr His
Tyr Ala Glu Ser Val Lys Gly Lys Phe Thr Ile Ser Arg Asp Asp Ser
Arg Ser Arg Leu Tyr Leu Gln Met Asn Asn Leu Arg Pro Glu Asp Ser
            100
Gly Ile Tyr Tyr Cys Thr Asp Gly Leu Gly Arg Pro Asn Trp Gly Gln
                            120
Gly Thr Leu Val Thr Val Ser Ala Ala Lys Thr Thr Pro Pro Ser Val
   130
                        135
                                            140
Tyr Pro Leu Ala Pro Cys Val
```

145

150

```
<210> 16
 <211> 15
 <212> PRT
 <213> Homo sapiens
 <400> 1.6
 Thr Ala Arg Ile Arg Ala Val Gly Leu Leu Thr Val Ile Ser Lys
<210> 17
 <211> 12
<212> PRT
<213> Homo sapiens
<400> 17
Val Asp Asp Ser Gln Asp Tyr Tyr Val Gly Lys Lys
<210> 18
<211> 15
<212> PRT
<213> Homo sapiens
<400> 18
Ser Leu Asn Cys Val Asp Asp Ser Gln Asp Tyr Tyr Val Gly Lys
<210> 19
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: RT-PCR PRIMER
<400> 19
tgcttgccct gttgatggca g
                                                                    21
<210> 20
<211> 22
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: RT-PCR PRIMER
<400> 20
ccagagcagc aggccgagtg ca
                                                                    22
<210> 21
<211> 25
<212> DNA
<213> Artificial Sequence
```

```
<220>
 <223> Description of Artificial Sequence: RT-PCR PRIMER
 <400> 21
 gggaattcgc acagccttca gggtc
                                                                     25
 <210> 22
 <211> 32-
 <212> DNA
 <213> Artificial Sequence
 <223> Description of Artificial Sequence: RT-PCR PRIMER
 <400> 22
 ggagaattca tggcactgcc ctgctgtgct ac
                                                                     32
 <210> 23
 <211> 30
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: RT-PCR PRIMER
 <400> 23
 ggagaattcc taatgggccc cgctggcgtt
                                                                     30
 <210> 24
 <211> 26
 <212> DNA
 <213> Artificial Sequence
 <220>
 <223> Description of Artificial Sequence: RT-PCR PRIMER
 <400> 24
 gggaagettg cacageette agggte
                                                                     26
 <210> 25
 <211> 39
 <212> DNA
<213> Artificial Sequence
<220>
 <223> Description of Artificial Sequence: RT-PCR PRIMER
<220>
<221> misc feature
<222> (18)
<223> a or g
<220>
<221> misc_feature
<222> (22)
<223> g or c
```

```
<220>
 <221> misc_feature
 <222> (28)
<223> g or t
<220>
<221> misc_feature
<222> (31)
<223> a or c
<220>
<221> misc_feature
<222> (34)
<223> g or c
<400> 25
ggcgatatcc accatggrat gsagctgkgt matsctctt
                                                                     39
<210> 26
<211> 39
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: RT-PCR PRIMER
<220>
<221> misc_feature
<222> (11)
<223> c or t
<220>
<221> misc_feature
<222> (25)..(26)
<223> a or g
<400> 26
agggaattca yctccacaca caggrrccag tggatagac
                                                                    39
<210> 27
<211> 39
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: RT-PCR PRIMER
<220>
<221> misc feature
<222> (17)
<223> a or g
<220>
<221> misc_feature
<222> (26)
<223> c or t
```

<220> <221> misc_feature <222> (33) <223> g or t

<400> 27 ggggatatcc accatgract tcgggytgag ctkggtttt

39